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Identification of methylation markers discriminating between CRC and normal blood

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An abbreviated version of this protocol was published in Science Translational Medicine in Dec 2019

Circulating tumor DNA methylation profiles enable early diagnosis prognosis prediction and screening for colorectal cancer

DOI: 10.1126/scitranslmed.aax7533

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The protocol for marker selection with omic dataset.docx

How to cite:(Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Luo, H. and Xu, R. (2020). Identification of methylation markers discriminating between CRC and normal blood. Bio-protocol Preprint. bio-protocol.org/prep267.
2. Luo, H., Zhao, Q., Wei, W., Zheng, L., Yi, S., Li, G., Wang, W., Sheng, H., Pu, H., Mo, H., Zuo, Z., Liu, Z., Li, C., Xie, C., Zeng, Z., Li, W., Hao, X., Liu, Y., Cao, S., Liu, W., Gibson, S., Zhang, K., Xu, G. and Xu, R.(2019). Circulating tumor DNA methylation profiles enable early diagnosis prognosis prediction and screening for colorectal cancer . Science Translational Medicine 12(524). DOI: 10.1126/scitranslmed.aax7533

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